1652

RAW SEQUENCE LISTING   DATE: 04/18/2001
PATENT APPLICATION: US/09/218,913B   TIME: 10:40:11

Input Set : A:\txt.98,736.Seq.Lst.2nd.Rev.txt
Output Set: N:\CRF3\04182001\I218913B.raw

ENTERED

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 3 <110> APPLICANT: Hall, Roderick L
 4        Poll, Christopher T.
 5        Newton, Benjamin B.
 6        Taylor, William J.A.
 8 <120> TITLE OF INVENTION: A Method for Accelerating the Rate of Mucociliary Clearance
 9 <130> FILE REFERENCE: 98,736
11 <140> CURRENT APPLICATION NUMBER: 09/218,913B
12 <141> CURRENT FILING DATE: 1998-12-22
14 <160> NUMBER OF SEQ ID NOS: 71
16 <170> SOFTWARE: Microsoft Word 97
18 <210> SEQ ID NO: 1
19 <211> LENGTH: 179
20 <212> TYPE: PRT
21 <213> ORGANISM: Homo sapien
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27 Val Gly Arg Cys Arg Ala Ser Met Pro Arg Trp Trp Tyr Asn Val Thr
28              20                  25                  30
30 Asp Gly Ser Cys Gln Leu Phe Val Tyr Gly Gly Cys Asp Gly Asn Ser
31          35                  40                  45
33 Asn Asn Tyr Leu Thr Lys Glu Glu Cys Leu Lys Lys Cys Ala Thr Val
34      50                  55                  60
36 Thr Glu Asn Ala Thr Gly Asp Leu Ala Thr Ser Arg Asn Ala Ala Asp
37 65                  70                  75                  80
39 Ser Ser Val Pro Ser Ala Pro Arg Arg Gln Asp Ser Glu Asp His Ser
40              85                  90                  95
42 Ser Asp Met Phe Asn Tyr Glu Glu Tyr Cys Thr Ala Asn Ala Val Thr
43              100                 105                 110
45 Gly Pro Cys Arg Ala Ser Phe Pro Arg Trp Tyr Phe Asp Val Glu Arg
46          115                 120                 125
48 Asn Ser Cys Asn Asn Phe Ile Tyr Gly Gly Cys Arg Gly Asn Lys Asn
49      130                 135                 140
51 Ser Tyr Arg Ser Glu Glu Ala Cys Met Leu Arg Cys Phe Arg Gln Gln
52 145                 150                 155                 160
54 Glu Asn Pro Pro Leu Pro Leu Gly Ser Lys Val Val Val Leu Ala Gly
55              165                 170                 175
57 Ala Val Ser
60 <210> SEQ ID NO: 2
61 <211> LENGTH: 197
62 <212> TYPE: PRT
63 <213> ORGANISM: Homo sapien
65 <220> FEATURE:
66 <221> NAME/KEY: sig_peptide
67 <222> LOCATION: 1..18
69 <400> SEQUENCE: 2
70 Ala Gly Ser Phe Leu Ala Trp Leu Gly Ser Leu Leu Leu Ser Gly Val
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Input Set : A:\txt.98,736.Seq.Lst.2nd.Rev.txt
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71  1                    5                    10                   15
73  Leu Ala Ala Asp Arg Glu Arg Ser Ile His Asp Phe Cys Leu Val Ser
74              20                   25                   30
76  Lys Val Val Gly Arg Cys Arg Ala Ser Met Pro Arg Trp Trp Tyr Asn
77          35                   40                   45
79  Val Thr Asp Gly Ser Cys Gln Leu Phe Val Tyr Gly Gly Cys Asp Gly
80      50                   55                   60
82  Asn Ser Asn Asn Tyr Leu Thr Lys Glu Glu Cys Leu Lys Lys Cys Ala
83  65                   70                   75                   80
85  Thr Val Thr Glu Asn Ala Thr Gly Asp Leu Ala Thr Ser Arg Asn Ala
86                   85                   90                   95
88  Ala Asp Ser Ser Val Pro Ser Ala Pro Arg Arg Gln Asp Ser Glu Asp
89              100                  105                  110
91  His Ser Ser Asp Met Phe Asn Tyr Glu Glu Tyr Cys Thr Ala Asn Ala
92          115                  120                  125
94  Val Thr Gly Pro Cys Arg Ala Ser Phe Pro Arg Trp Tyr Phe Asp Val
95      130                  135                  140
97  Glu Arg Asn Ser Cys Asn Asn Phe Ile Tyr Gly Gly Cys Arg Gly Asn
98  145                  150                  155                  160
100 Lys Asn Ser Tyr Arg Ser Glu Glu Ala Cys Met Leu Arg Cys Phe Arg
101                  165                  170                  175
103 Gln Gln Glu Asn Pro Pro Leu Pro Leu Gly Ser Lys Val Val Val Leu
104              180                  185                  190
106 Ala Gly Ala Val Ser
107          195
109 <210> SEQ ID NO: 3
110 <211> LENGTH: 153
111 <212> TYPE: PRT
112 <213> ORGANISM: Homo sapien
114 <400> SEQUENCE: 3
115 Ile His Asp Phe Cys Leu Val Ser Lys Val Val Gly Arg Cys Arg Ala
116 1                    5                    10                   15
118 Ser Met Pro Arg Trp Trp Tyr Asn Val Thr Asp Gly Ser Cys Gln Leu
119              20                   25                   30
121 Phe Val Tyr Gly Gly Cys Asp Gly Asn Ser Asn Asn Tyr Leu Thr Lys
122          35                   40                   45
124 Glu Glu Cys Leu Lys Lys Cys Ala Thr Val Thr Glu Asn Ala Thr Gly
125      50                   55                   60
127 Asp Leu Ala Thr Ser Arg Asn Ala Ala Asp Ser Ser Val Pro Ser Ala
128 65                   70                   75                   80
130 Pro Arg Arg Gln Asp Ser Glu Asp His Ser Ser Asp Met Phe Asn Tyr
131                  85                   90                   95
133 Glu Glu Tyr Cys Thr Ala Asn Ala Val Thr Gly Pro Cys Arg Ala Ser
134          100                  105                  110
136 Phe Pro Arg Trp Tyr Phe Asp Val Glu Arg Asn Ser Cys Asn Asn Phe
137          115                  120                  125
139 Ile Tyr Gly Gly Cys Arg Gly Asn Lys Asn Ser Tyr Arg Ser Glu Glu
140          130                  135                  140
142 Ala Cys Met Leu Arg Cys Phe Arg Gln
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            Input Set : A:\txt.98,736.Seq.Lst.2nd.Rev.txt
            Output Set: N:\CRF3\04182001\I218913B.raw

143 145   150
145 <210> SEQ ID NO: 4
146 <211> LENGTH: 58
147 <212> TYPE: PRT
148 <213> ORGANISM: Homo sapien
150 <400> SEQUENCE: 4
151 Ile His Asp Phe Cys Leu Val Ser Lys Val Val Gly Arg Cys Arg Ala
152 1   5   10   15
154 Ser Met Pro Arg Trp Trp Tyr Asn Val Thr Asp Gly Ser Cys Gln Leu
155   20   25   30
157 Phe Val Tyr Gly Gly Cys Asp Gly Asn Ser Asn Asn Tyr Leu Thr Lys
158   35   40   45
160 Glu Glu Cys Leu Lys Lys Cys Ala Thr Val
161   50   55
163 <210> SEQ ID NO: 5
164 <211> LENGTH: 51
165 <212> TYPE: PRT
166 <213> ORGANISM: Homo sapien
168 <400> SEQUENCE: 5
169 Cys Leu Val Ser Lys Val Val Gly Arg Cys Arg Ala Ser Met Pro Arg
170 1   5   10   15
172 Trp Trp Tyr Asn Val Thr Asp Gly Ser Cys Gln Leu Phe Val Tyr Gly
173   20   25   30
175 Gly Cys Asp Gly Asn Ser Asn Asn Tyr Leu Thr Lys Glu Glu Cys Leu
176   35   40   45
178 Lys Lys Cys
179   50
181 <210> SEQ ID NO: 6
182 <211> LENGTH: 58
183 <212> TYPE: PRT
184 <213> ORGANISM: Homo sapien
186 <400> SEQUENCE: 6
187 Tyr Glu Glu Tyr Cys Thr Ala Asn Ala Val Thr Gly Pro Cys Arg Ala
188 1   5   10   15
190 Ser Phe Pro Arg Trp Tyr Phe Asp Val Glu Arg Asn Ser Cys Asn Asn
191   20   25   30
193 Phe Ile Tyr Gly Gly Cys Arg Gly Asn Lys Asn Ser Tyr Arg Ser Glu
194   35   40   45
196 Glu Ala Cys Met Leu Arg Cys Phe Arg Gln
197   50   55
199 <210> SEQ ID NO: 7
200 <211> LENGTH: 51
201 <212> TYPE: PRT
202 <213> ORGANISM: Homo sapien
204 <400> SEQUENCE: 7
205 Cys Thr Ala Asn Ala Val Thr Gly Pro Cys Arg Ala Ser Phe Pro Arg
206 1   5   10   15
208 Trp Tyr Phe Asp Val Glu Arg Asn Ser Cys Asn Asn Phe Ile Tyr Gly
209   20   25   30

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211 Gly Cys Arg Gly Asn Lys Asn Ser Tyr Arg Ser Glu Glu Ala Cys Met
212         35              40              45
214 Leu Arg Cys
215     50
217 <210> SEQ ID NO: 8
218 <211> LENGTH: 92
219 <212> TYPE: PRT
220 <213> ORGANISM: Homo sapien
222 <400> SEQUENCE: 8
223 Ala Asp Arg Glu Arg Ser Ile His Asp Phe Cys Leu Val Ser Lys Val
224 1               5               10              15
226 Val Gly Arg Cys Arg Ala Ser Met Pro Arg Trp Trp Tyr Asn Val Thr
227         20              25              30
229 Asp Gly Ser Cys Gln Leu Phe Val Tyr Gly Gly Cys Asp Gly Asn Ser
230         35              40              45
232 Asn Asn Tyr Leu Thr Lys Glu Glu Cys Leu Lys Lys Cys Ala Thr Val
233     50              55              60
235 Thr Glu Asn Ala Thr Gly Asp Leu Ala Thr Ser Arg Asn Ala Ala Asp
236 65              70              75              80
238 Ser Ser Val Pro Ser Ala Pro Arg Arg Gln Asp Ser
239             85              90
241 <210> SEQ ID NO: 9
242 <211> LENGTH: 708
243 <212> TYPE: DNA
244 <213> ORGANISM: Homo sapien
246 <220> FEATURE:
247 <221> NAME/KEY: misc_feature
248 <222> LOCATION: 679..708
249 <223> OTHER INFORMATION: /note= "n at positions 622, 679, 707 is any nucleic acid"
251 <400> SEQUENCE: 9
252 ggccgggtcg tttctcgcct ggctgggatc gctgctcctc tctggggtcc tggcggccga      60
254 ccgagaacgc agcatccacg acttctgcct ggtgtcgaag gtggtgggca gatgccgggc     120
256 ctccatgcct aggtggtggt acaatgtcac tgacggatcc tgccagctgt ttgtgtatgg     180
258 gggctgtgac ggaaacagca ataattacct gaccaaggag gagtgcctca agaaatgtgc     240
260 cactgtcaca gagaatgcca cgggtgacct ggccaccagc aggaatgcag cggattcctc     300
262 tgtcccaagt gctcccagaa ggcaggattc tgaagaccac tccagcgata tgttcaacta     360
264 tgaagaatac tgcaccgcca acgcagtcac tgggccttgc cgtgcatcct cccacgctg      420
266 gtactttgac gtggagagga actcctgcaa taacttcatc tatggaggct gccggggcaa     480
268 taagaacagc taccgctctg aggaggcctg catgctccgc tgcttccgcc agcaggagaa     540
270 tcctcccctg ccccttggct caaaggtggt ggttctggcc ggggctgttt cgtgatggtg     600
272 ttgatccttt tcctggggag cntccatggt cttactgatt ccgggtggca aggaggaacc     660
274 aggagcgtgc cctgcgganc gtctggagct tcggagatga caagggnt                 708
276 <210> SEQ ID NO: 10
277 <211> LENGTH: 235
278 <212> TYPE: PRT
279 <213> ORGANISM: Homo sapien
281 <220> FEATURE:
282 <221> NAME/KEY: peptide
283 <222> LOCATION: 1..235
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RAW SEQUENCE LISTING   DATE: 04/18/2001
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Input Set : A:\txt.98,736.Seq.Lst.2nd.Rev.txt
Output Set: N:\CRF3\04182001\I218913B.raw

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284 <223> OTHER INFORMATION: /note= "Xaa at positions 198, 201, 226, and 233 are unknown
285       amino acids"
287 <400> SEQUENCE: 10
288 Ala Gly Ser Phe Leu Ala Trp Leu Gly Ser Leu Leu Leu Ser Gly Val
289 1               5                   10                  15
291 Leu Ala Ala Asp Arg Glu Arg Ser Ile His Asp Phe Cys Leu Val Ser
292             20                  25                  30
294 Lys Val Val Gly Arg Cys Arg Ala Ser Met Pro Arg Trp Trp Tyr Asn
295         35                  40                  45
297 Val Thr Asp Gly Ser Cys Gln Leu Phe Val Tyr Gly Gly Cys Asp Gly
298     50                  55                  60
300 Asn Ser Asn Asn Tyr Leu Thr Lys Glu Glu Cys Leu Lys Lys Cys Ala
301 65                  70                  75                  80
303 Thr Val Thr Glu Asn Ala Thr Gly Asp Leu Ala Thr Ser Arg Asn Ala
304             85                  90                  95
306 Ala Asp Ser Ser Val Pro Ser Ala Pro Arg Arg Gln Asp Ser Glu Asp
307             100                 105                 110
309 His Ser Ser Asp Met Phe Asn Tyr Glu Glu Tyr Cys Thr Ala Asn Ala
310         115                 120                 125
312 Val Thr Gly Pro Cys Arg Ala Ser Phe Pro Arg Trp Tyr Phe Asp Val
313     130                 135                 140
315 Glu Arg Asn Ser Cys Asn Asn Phe Ile Tyr Gly Gly Cys Arg Gly Asn
316 145                 150                 155                 160
318 Lys Asn Ser Tyr Arg Ser Glu Glu Ala Cys Met Leu Arg Cys Phe Arg
319             165                 170                 175
321 Gln Gln Glu Asn Pro Pro Leu Pro Leu Gly Ser Lys Val Val Val Leu
322             180                 185                 190
324 Ala Gly Ala Val Ser Xaa Trp Cys Xaa Ser Phe Ser Trp Gly Ala Ser
325         195                 200                 205
327 Met Val Leu Leu Ile Pro Gly Gly Lys Glu Glu Pro Gly Ala Cys Pro
328     210                 215                 220
330 Ala Xaa Arg Leu Glu Leu Arg Arg Xaa Gln Gly
331 225                 230                 235
333 <210> SEQ ID NO: 11
334 <211> LENGTH: 179
335 <212> TYPE: PRT
336 <213> ORGANISM: Homo sapien
338 <220> FEATURE:
339 <221> NAME/KEY: peptide
340 <222> LOCATION: 1..170
341 <223> OTHER INFORMATION: /note= "Xaa at positions 8, 17, 19, 21-26, 40, 42, 45-47, 52, 64,
342       103, 112, 114, 116-121, 135, 137, 140-142, 147, and 159 is any
343       amino acid residue"
345 <400> SEQUENCE: 11
346 Ala Asp Arg Glu Arg Ser Ile Xaa Asp Phe Cys Leu Val Ser Lys Val
347 1               5                   10                  15
349 Xaa Gly Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Trp Trp Tyr Asn Val Thr
350         20                  25                  30
352 Asp Gly Ser Cys Gln Leu Phe Xaa Tyr Xaa Gly Cys Xaa Xaa Xaa Ser
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W-->
W-->
W-->
W-->
W-->

file://

4/18/01

VERIFICATION SUMMARY   DATE: 04/18/2001
PATENT APPLICATION: US/09/218,913B   TIME: 10:40:12

Input Set : A:\txt.98,736.Seq.Lst.2nd.Rev.txt
Output Set: N:\CRF3\04182001\I218913B.raw

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L:9 M:283 W: Missing Blank Line separator, <130> field identifier
L:272 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:274 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:324 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10
L:330 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10
L:346 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11
L:349 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11
L:352 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11
L:355 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11
L:364 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11
L:367 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11
L:370 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11
L:373 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11
L:419 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12
L:427 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:13
L:456 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:459 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:462 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:490 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14
L:492 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14
L:505 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15
L:508 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15
L:529 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15
L:532 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15
L:535 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15
L:549 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16
L:563 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16
L:576 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
L:586 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
L:588 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
L:598 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:18
L:992 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:45
L:1099 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:47
L:1206 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:49
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